

1
 GAATTCCGAT TTAGCCTCAT ACTGCTTCTC ACATTACATT GGGATGCGCT
 51
 TTGCAAACAC ACCCCAATGC TGCCTCATT GGGGAAGAGG TTGCTGCGAA
 101
 GAAGCAAACC CTTAAGAACG TCACAACTA CATTACTGAT ATCATCTGCA
 151
 AGCGTGCAGA TCTTGTTAC AACTATGGGG TTATCCTTAT ACCAGAAGGC
 201
 CTGATTGATT TCATCCCAGA GGTTCAAAA CTCATCGCAG AATTGAATGA
 251
 AATTTTGGCA CATGATGTGG TTGATGAGGC AGGGGCCTGG AAAAGCAAGC
 301
 TTCAGCCTGA ATCAAAGGAG CTGTTTGAGT TTTTGCCCA AACTATTCAG
 351
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 401
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 451
 AGAGAAAAGC AGAGGGGAGA TACTCTGCAC ATTCAGAGG GCAAGCTCAT
 501
 TTCTTTGGGT ACGAAGGAAG ATGTGGCCTT CCTACCAATT TTGATTCTAA
 551
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 651
 TGGACTGTTG GTGGAACAGC ATTGACATCA CTGATGGATG TTGAGAGGAG
 701
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 751
 CTGCACCTTT CAAGAAATAT GCATCAATGC GGGATGAGTG GGCCACCAAG
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 851
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 901
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 1001
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 1051
 AGAACTGGTT TTAGCATTTT TTGTATGATT TACGCACCAA CTGACTTGTC
 1101
 TTGTAACCCT GATTCTGTTG CACTGGTTGC /ATCTCGTGA GAATGAACAA
 1151
 GTTGATATGA GGCTAAATCG GAATTC

Figure 1.

1
 ATGGCGGCGC CGAGCGGACC ATCACCTGGG ACTGGGAGGT TGGCGTCGGT
 51
 TTACAGCGAG GTGCAGACGA GCCGCCTCCA TCACGCGATC CGGCTCCCCT
 101
 CCGTCCTCTG CTCCCAATTC TCCCTCGTCG ATGGACCTCC CAGCTCAGCC
 151
 ACGGGGAACC CGGATGAGAT CGCGAAGCTG TTCCCTAACT TGTTTGGGCA
 201
 GCCGTCGGCG ACATTGGTGC CGGCCAAGA GCGGTGGAG GGAAGGCGC
 251
 TGAAGGTCGG GGTGGTGCTC TCTGGTGGAC AAGCACCCGG TGGGCACAAT
 301
 GTGATCTGCG GTATCTTCGA TTTCTTGCAG AAACACGCAA AGGGAAGCAC
 351
 AATGTATGGA TTCAAAGGAG GCCCAGCAGG GGTGATGAAG TGCAAGTACG
 401
 TCAAAC TAA TACCGATTTC GTCTATCCCT ACAGAAACCA GGGTGGTTTT
 451
 GATATGATCT GTAGTGGAAG GGATAAGATT GAAACACCAG AGCAGTTTAA
 501
 GCAAGCCGAA GATACAGCCA ACAAACTTGA GTTGGACGGA CTTGTTGTTA
 551
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 601
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 651
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 701
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 751
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 TCACATTACA TTGGGATGCG CTTTGCAAAC ACACCCCAAT GCTGCACTCA
 851
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 901
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 951
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 1001
 ATATCATTGC TGAATTGAAT GAAATTTTGG CACATGATGT TGTTGATGAG
 1051
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 1101
 GTTTTGGCC AAAACTATTC AGGAGCAACT TATGCTTGAA AGGGGCCCCC
 1151
 ATGGCAATGT TCAGGTTGCA AAAATTGAA CCGAGAAAAT GCTTATTAGC
 1201
 ATGGTGGAAA CTGAACTGGA GAAGAGAAA GCAGAGGGGA GATACTCTGC

Figure 2

1251
ACATTTTCAGA GGGCAAGCTC ATTTCTTTGG GTACGAAGGA AGATGTGGCC
1301
TTCCTACCAA TTTTGATTCT AACTATTGCT ATGCATTAGG CTATGGGGCT
1351
GGTGCCCTTC TCCAAAGTGG GAAGACAGGA CTTATTTTCAT CGGTTGGCAA
1401
CCTTGCGGCT CCAGTAGAAG AATGGACTGT TGGTGGGAACA GCATTGACAT
1451
CACTGATGGA TGTGGAGAGG AGGCATGGCA AGTTCAAGCC AGTGATCGAG

1501
AAGGCTATGG TGGAACCTGA TGCTGCACCT TTCAAGAAAT ATGCATCAAT
1551
GCGGGATGAG TGGGCCACCA AGAACAGATA CATCAGCCCT GGCCCCATCC
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1651
CTCGGTGCTG AGTTATAG

Figure 2 cont.

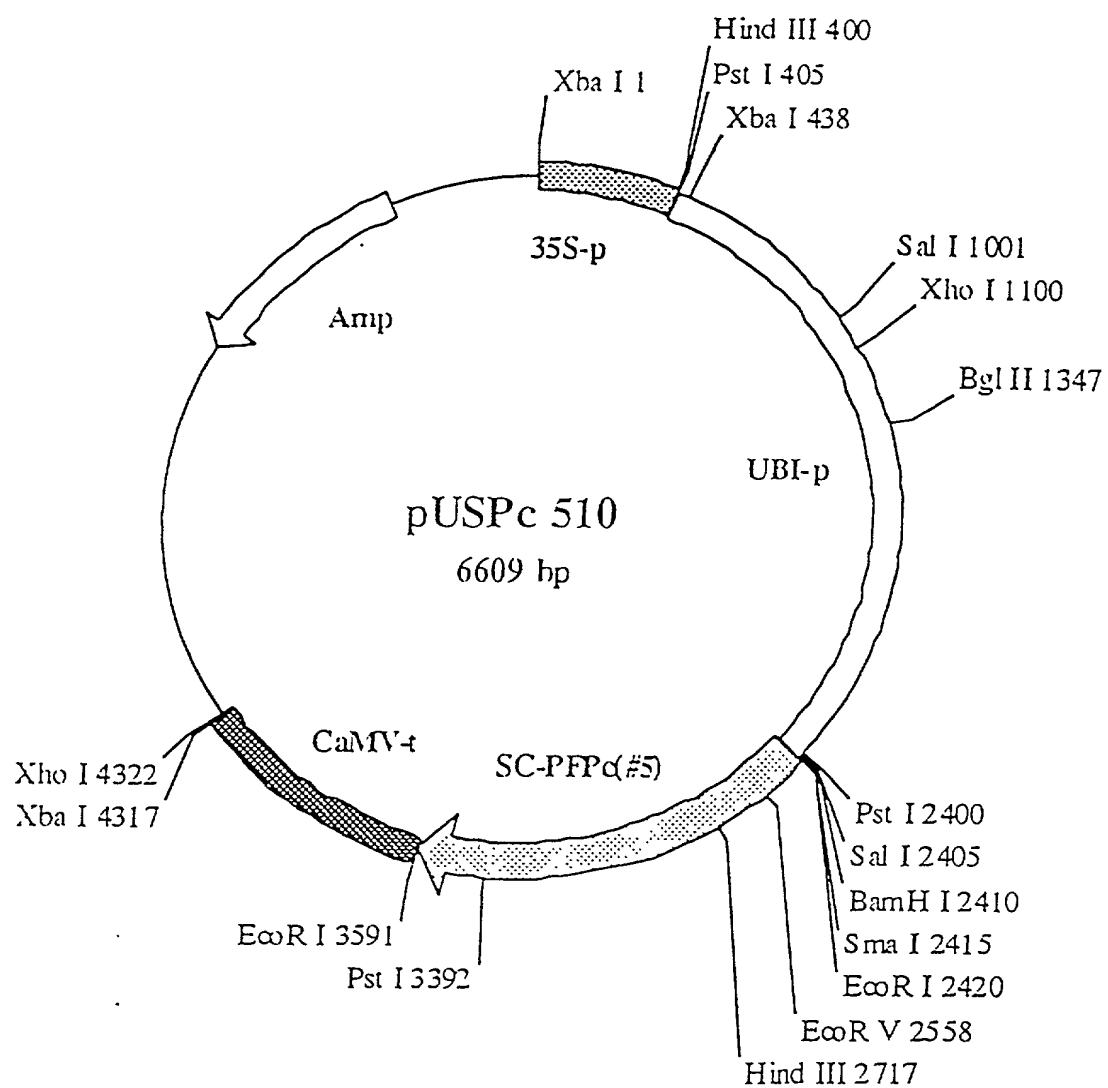


Figure 3

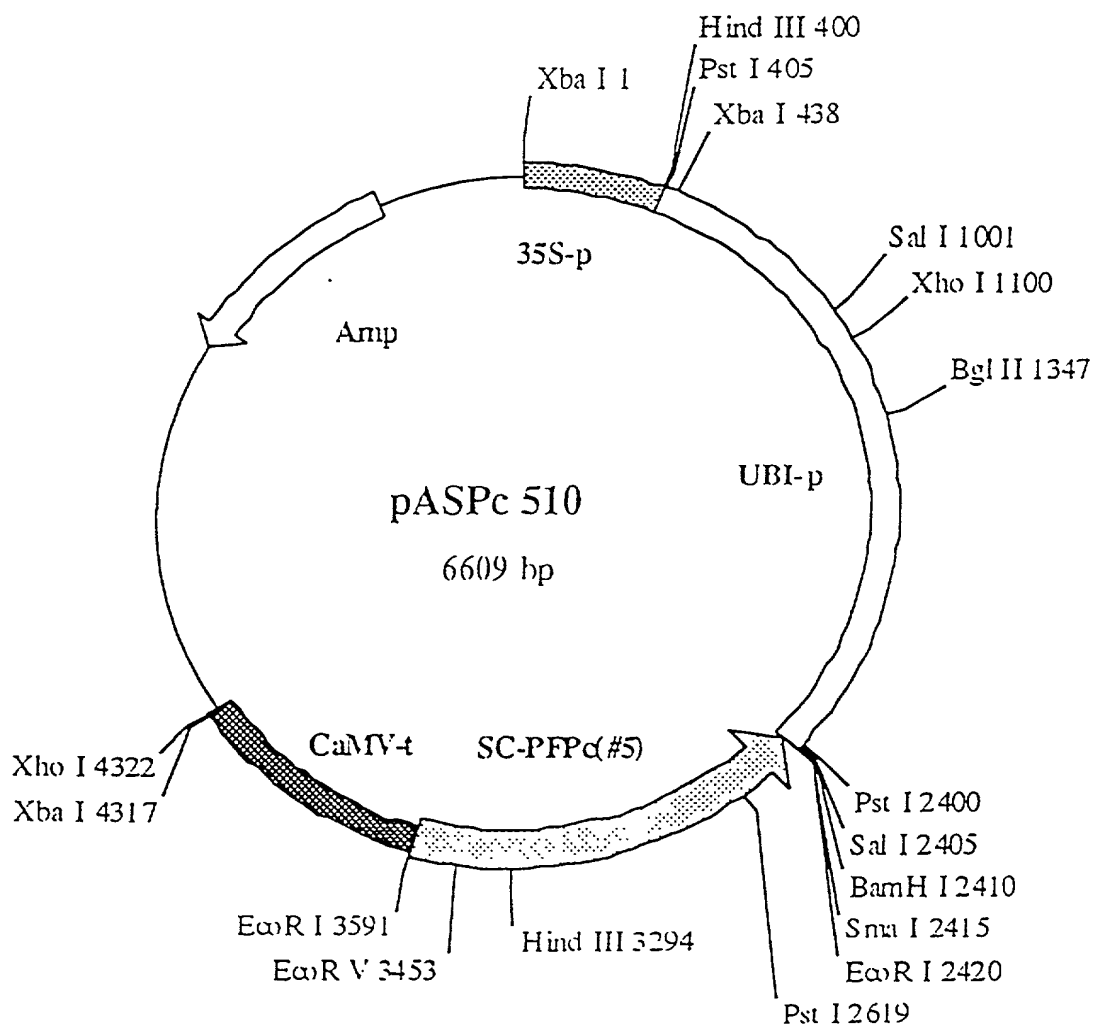


Figure 4

1 2 3 4 5 6 7 8 9 10 11 12



Figure 5.

1 2 3 4 5

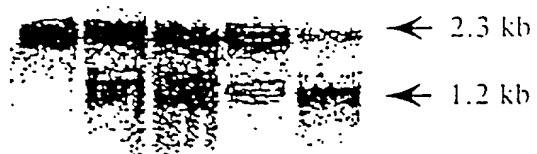


Figure 6

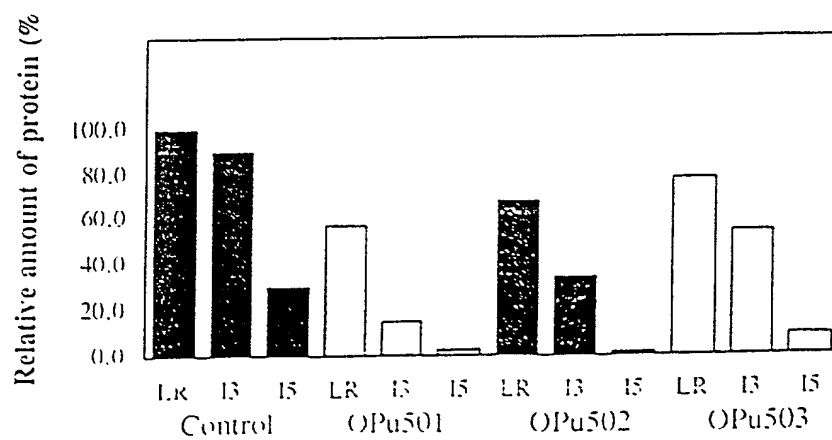


Figure 7